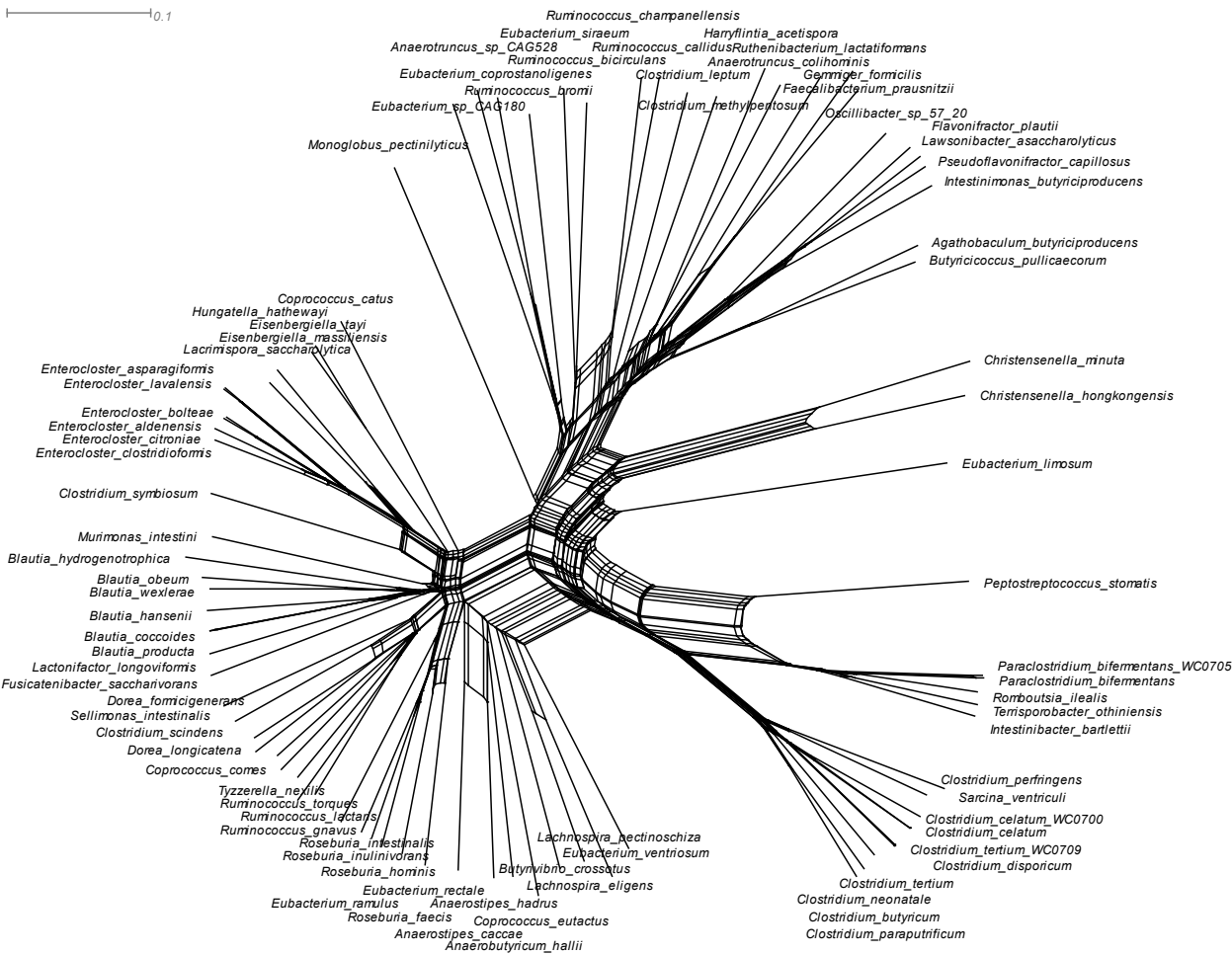


Supplementary Figure 3: Split decomposition of phylogenetic trees base on core genes alignment (A) and AAI (B). The analysis was performed utilizing gene alignments for core genome genes and distance matrix for AAI.

A



B

